FIG. 1

GCTGCTGCTGCTGCTGCTCGTGCCCGTGCCGCTGCTGCCGCTGCTGGCCCAAGGGCCCGGAGGGGCGC
TGGGAAACCGGCATGCGGTGTACTGGAACAGCTCCAACCAGCACCTGCGGCGAGAGGGCTACACCGTGCAG
GTGAACGTGAACGACTATCTGGATATTTACTGCCCGCACTACAACAGCTCGGGGGTGGGCCCCGGGGCGGG
ACCGGGGCCCGGAGGCGGGGCAGAGCAGTACGTGCTGTACATGGTGAGCCGCAACGGCTACCGCACCTGCA
ACGCCAGCCAGGGCTTCAAGCGCTGGGAGTGCAACCGGCCGCACGCCCCGCACAGCCCCATCAAGTTCTCG
GAGAAGTTCCAGCGCTACAGCGCCTTCTCTCTGGGCTACGAGTTCCACGCCGGCCACGAGTACTACTACAT
CTCCACGCCCACTCACAACCTGCACTGGAAGTGTCTGAGGATGAAGGTGTTCGTCTGCTGCGCCTCCACAT
CGCACTCCGGGGAGAAGCCGGTCCCCACTCTCCCCCAGTTCACCATGGGCCCCAATGTGAAGATCAACGTG
CTGGAAGACTTTGAGGGAGAGAACCCTCAGGTGCCCAAGCTTGAGAAGAGCATCAGCGGGACCAGCCCCAA
ACGGGAACACCTGCCCCTGGCCGTGGGCATCGCCTTCTTCCTCATGACGTTCTTGGCCTCCTAGCTCTGCC
CCCTCCCCTGGGGGGGGAGAGATGGGGCGGGGCTTGGAAGGAGCAGGGAGCCTTTGGCCTCTCCAAGGGAA
GCCTAGTGGGCCTAGACCCCTCCTCCCATGGCTAGAAGTGGGGCCTGCACCATACATCTGTGTCCGCCCCC
TCTACCCCTTCCCCCCACGTAGGGCACTGTAGTGGACCAAGCACGGGGACAGCCATGGGTCCCGGGCGGCC
TTGTGGCTCTGGTAATGTTTGGTACCAAACTTGGGGGCCAAAAAGGGCAGTGCTCAGGACTCCCTGGCCCC
TGGTACCTTTCCCTGACTCCTGGTGCCCTCTCCCTTTGTCCCCCCAGAGAGACATATGCCCCCAGAGAGAG
CAAATCGAAGCGTGGGAGGCACCCCCATTGCTCTCCTCCAGGGGCAGAACATGGGGAGGGGACTAGATGGG
CAAGGGGCAGCACTGCCTGCTGCTTCCTTCCCCTGTTTACAGCAATAAGCACGTCCTCCTCCCCCACTCCC
ACTTCCAGGATTGTGGTTTGGATTGAAACCAAGTTTACAAGTAGACACCCCTGGGGGGGGCGGGCAGTGGAC
AAGGATGCCAAGGGGTGGGCATTGGGGTGCCAGGCAGGCATGTACAGACTCTATATCTCTATATATAATGT
ACAGACAGACAGAGTCCCTTCCCTCTTTAACCCCCTGACCTTTCTTGACTTCCCCTTCAGCTTCAGACCCC
TTCCCCACCAGGCTAGGCCCCCCACACCTGGGGGACCCCCTGGCCCCTCTTTTGTCTTCTGTGAAGACAGG
ACCTATGCAACGCACAGACACTTTTGGAGACCGTAAAACAACAGCGCCCCCTCCCTTCCAGCCCTGAGCCG
GGAACCATCTCCCAGGACCTTGCCCTGCTCACCCTATGTGGTCCCACCTATCCTCCTGGGCCTTTTTCAAG
TGCTTTGGCTGTGACTTTCATACTCTGCTCTTAGTCTAAAAAAAATAAACTGGAGATAAAAATAAAAAAAA
TACCTCGAGAAAAAAAAAAAAAAAAAAAAAAAAAAAA

FIG. 2

MAAAPLLLLLLLLVPVPLLPLLAQGPGGALGNRHAVYWNSSNQHLRREGYTVQVNVNDYLDIYCPHYNSSGV
GPGAGPGPGGGAEQYVLYMVSRNGYRTCNASQGFKRWECNRPHAPHSPIKFSEKFQRYSAFSLGYEFHAGH
EYYYISTPTHNLHWKCLRMKVFVCCASTSHSGEKPVPTLPQFTMGPNVKINVLEDFEGENPQVPKLEKSIS
GTSPKREHLPLAVGIAFFLMTFLAS

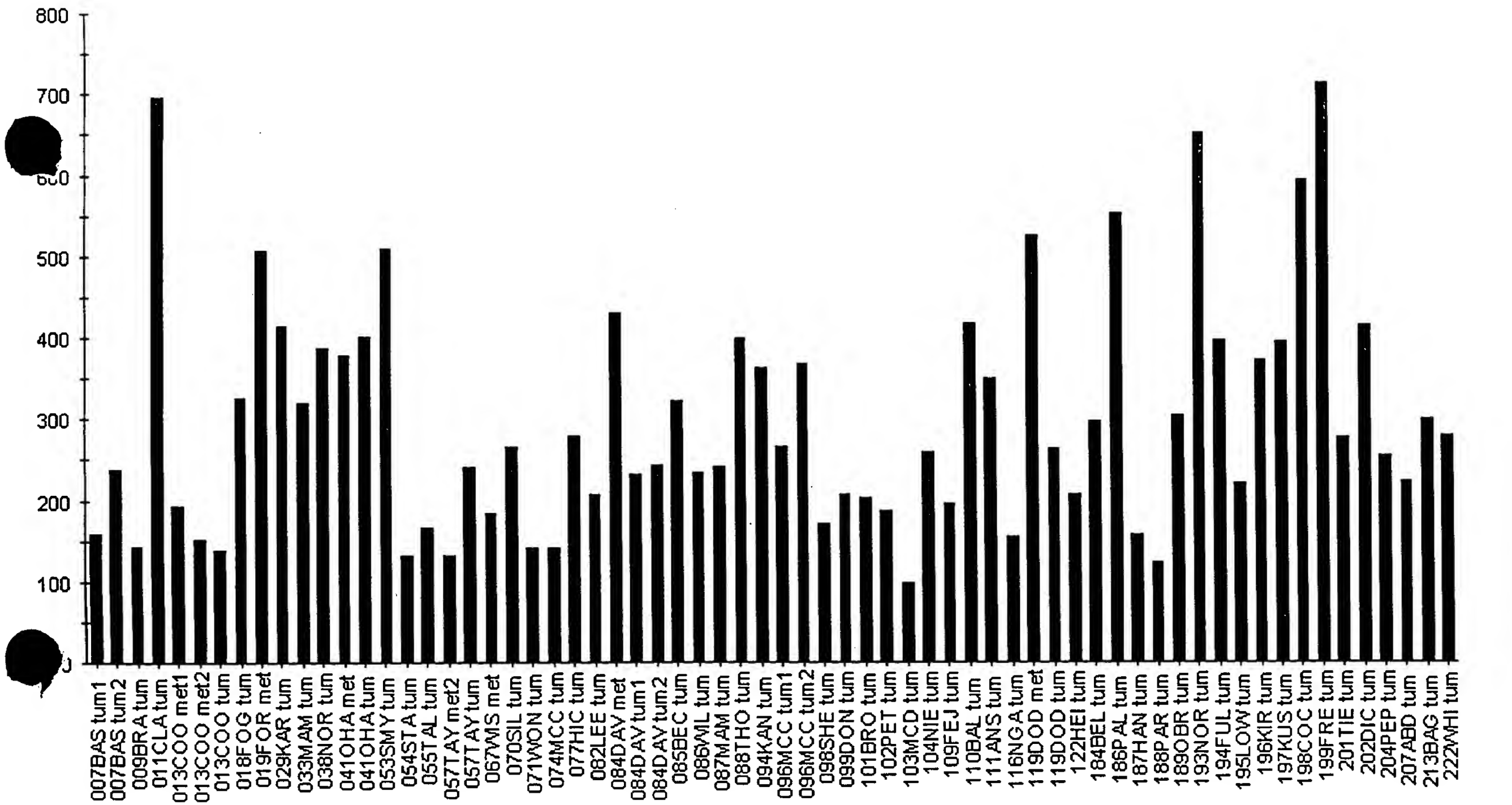

FIG. 3A

09733756 . 120800

09733756.120800

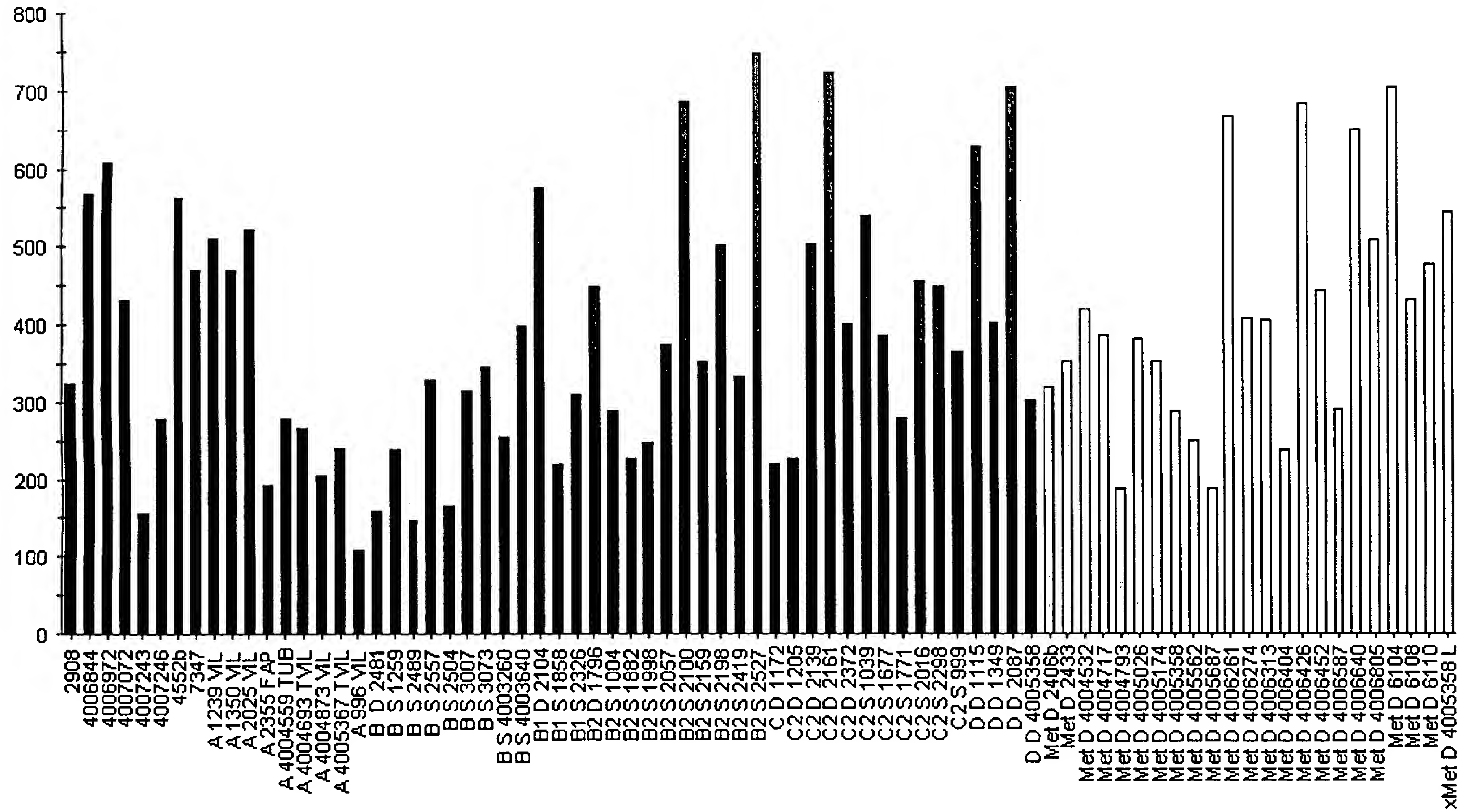

FIG. 3B

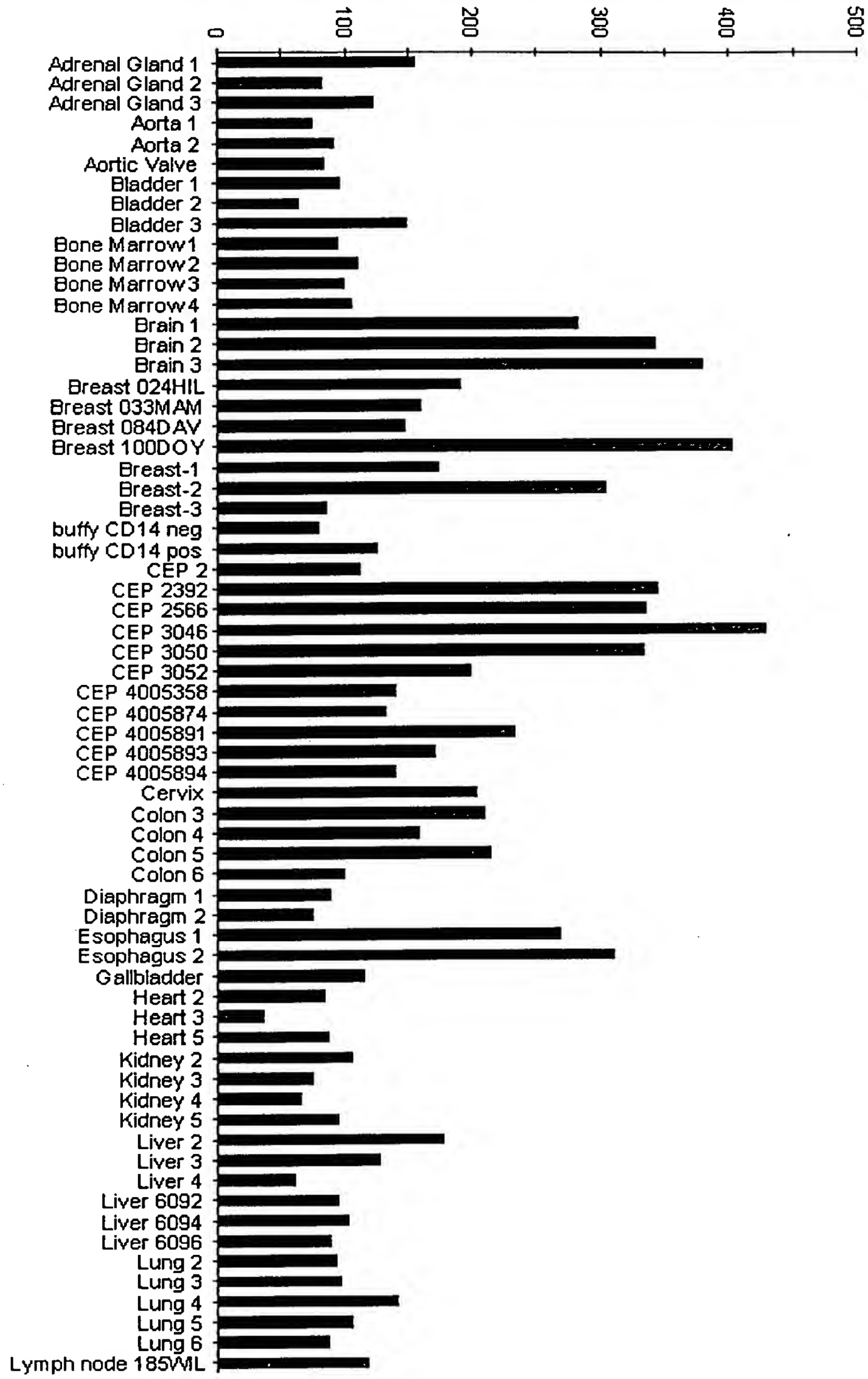
0
100
200
300
400
500
Adrenal Gland 1
Adrenal Gland 2
Adrenal Gland 3
Aorta 1
Aorta 2
Aortic Valve
Bladder 1
Bladder 2
Bladder 3
Bone Marrow1
Bone Marrow2
Bone Marrow3
Bone Marrow4
Brain 1
Brain 2
Brain 3
Breast 024HIL
Breast 033MAM
Breast 084DAV
Breast 100DOY
Breast-1
Breast-2
Breast-3
buffy CD14 neg
buffy CD14 pos
CEP 2
CEP 2392
CEP 2566
CEP 3046
CEP 3050
CEP 3052
CEP 4005358
CEP 4005874
CEP 4005891
CEP 4005893
CEP 4005894
Cervix
Colon 3
Colon 4
Colon 5
Colon 6
Diaphragm 1
Diaphragm 2
Esophagus 1
Esophagus 2
Gallbladder
Heart 2
Heart 3
Heart 5
Kidney 2
Kidney 3
Kidney 4
Kidney 5
Liver 2
Liver 3
Liver 4
Liver 6092
Liver 6094
Liver 6096
Lung 2
Lung 3
Lung 4
Lung 5
Lung 6
Lymph node 185WIL

FIG. 3C

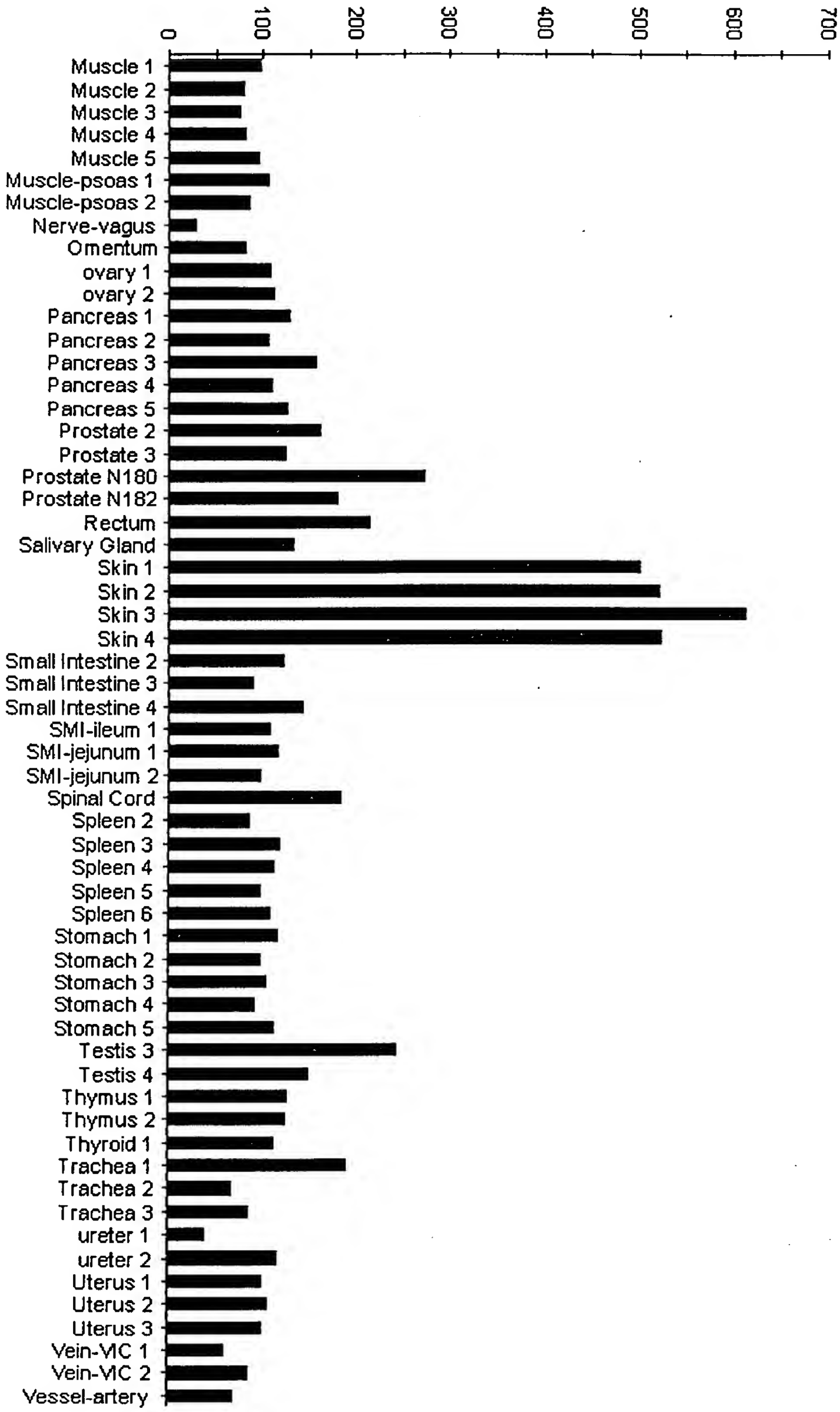

FIG. 3D

Doc Code: AP.PRE.REQ